## **Amendments to the Claims:**

This listing of claims will replace all prior versions, and listings of claims in the application:

## **Listing of Claims:**

1. - 79. (Canceled)

80. (Currently amended) A polypeptide consisting essentially of: a first and a second interactor domain, and a circularly permutated TEM-1 β-lactamase protein;

wherein the first interactor domain binds to a single ligand, and the first interactor domain is selected from the group consisting of an antibody, an antigen, a first monomer of a hetero-dimerizing helix, a second monomer of a hetero-dimerizing helix, a receptor, a member of an expressed sequence library, a member of a constrained peptide library, and a scaffold peptide member of a thioredoxin peptide library;

wherein the second interactor domain binds to said single ligand, and the second interactor domain is selected from the group consisting of an antibody, an antigen, a first monomer of a hetero-dimerizing helix, a second monomer of a hetero-dimerizing helix, a receptor, a member of an expressed sequence library, a member of a constrained peptide library, and a seaffold peptide member of a thioredoxin peptide library;

wherein the first interactor domain is fused to the circularly permutated  $\beta$ -lactamase protein through the N-terminal break-point of the circularly permutated  $\beta$ -lactamase protein and the second interactor domain is fused to the circularly permutated  $\beta$ -lactamase protein through the C-terminal break-point of the circularly permutated  $\beta$ -lactamase protein,

wherein said N-terminal break-point and said C-terminal break-point of the circularly permutated TEM-1 β-lactamase protein are within 10 amino acids in either direction from a junction of between 2 amino acid residues in a solvent exposed loop between amino acid

residues Thr 195 and Ala 202 and, located between alpha-helices 7 and 8 of said TEM-1 β-lactamase protein,

wherein said circularly permutated TEM-1 β-lactamase protein is functionally reconstituted only upon binding of said first interactor domain and to said second interactor domain to said single ligand.

81-84. (Canceled)

85. (Currently amended) The polypeptide of claim 80, wherein said circularly permutated  $\beta$ -lactamase protein consists of amino acids 26 to 288 of the following sequence prior to circular permutation

His Pro Glu Thr Leu Val Lys Val Lys Asp Ala Glu Asp Gln Leu Gly Ala Arg Val Gly Tyr Ile Glu Leu Asp Leu Asn Ser Gly Lys Ile Leu Glu Ser Phe Arg Pro Glu Glu Arg Phe Pro Met Met Ser Thr Phe Lys Val Leu Leu Cys Gly Ala Val Leu Ser Arg Ile Asp Ala Gly Gln Glu Gln Leu Gly Arg Arg Ile His Tyr Ser Gln Asn Asp Leu Val Glu Tyr Ser Pro Val Thr Glu Lys His Leu Thr Asp Gly Met Thr Val Arg Glu Leu Cys Ser Ala Ala Ile Thr Met Ser Asp Asn Thr Ala Ala Asn Leu Leu Leu Thr Thr Ile Gly Gly Pro Lys Glu Leu Thr Ala Phe Leu His Asn Met Gly Asp His Val Thr Arg Leu Asp Arg Trp Glu Pro Glu Leu 

Asn Glu Ala Ile Pro Asn Asp Glu Arg Asp Thr Thr Met Pro Val Ala 180 170 175 185 Met Ala Thr Thr Leu Arg Lys Leu Leu Thr Gly Glu Leu Leu Thr Leu 190 195 200 Ala Ser Arg Gln Gln Leu Ile Asp Trp Met Glu Ala Asp Lys Val Ala 205 210 215 Gly Pro Leu Leu Arg Ser Ala Leu Pro Ala Gly Trp Phe Ile Ala Asp 220 225 230 Lys Ser Gly Ala Gly Glu Arg Gly Ser Arg Gly Ile Ile Ala Ala Leu 235 240 245 Gly Pro Asp Gly Lys Pro Ser Arg Ile Val Val Ile Tyr Thr Thr Gly 250 255 260 265 Ser Gln Ala Thr Met Asp Glu Arg Asn Arg Gln Ile Ala Glu Ile Gly 270 275 280 Ala Ser Leu Ile Lys His Trp

20.5

285

(SEQ ID NO: 2);

wherein said C-terminal break-point and said N-terminal break-point is between amino acid residues Glu 197 and Leu 198.

86-89. (Canceled).

90. (Currently amended) The polypeptide of claim 80, wherein when said first interactor domain and said second interactor domain bind to a single ligand, and is an antibody, said second interactor domain is a first monomer of a hetero-dimerizing helix, and said ligand is an antigen-second monomer of hetero-dimerizing helix fusion protein, wherein the antibody specifically binds to the antigen; or

when wherein said circularly permutated TEM-1 β-lactamase protein is functionally reconstituted only upon binding of said first interactor domain is an antibody, and

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said second interactor domain is a first monomer of a hetero-dimerizing helix, and to said ligand is a second monomer of hetero-dimerizing helix-antigen fusion protein, wherein the antibody specifically binds to the antigen; or

when said first interactor domain is an antigen, said second interactor domain is a first monomer of a hetero-dimerizing helix, and said ligand is an antibody- second monomer of hetero-dimerizing helix fusion protein, wherein the antigen specifically binds to the antibody; or

when said first interactor domain is a first monomer of a hetero-dimerizing helix, said second interactor domain is an antigen, and said ligand is an antibody-second monomer of hetero-dimerizing helix fusion protein, wherein the antigen specifically binds to the antibody.

91. (Currently amended) The polypeptide of claim 90, wherein <u>said ligand is</u> comprised of an antigen fused to a second monomer of a hetero-dimerizing helix protein, said first <u>interactor domain is an antibody</u>, and said second interactor domain is a first monomer of a hetero-dimerizing helix, or

wherein said ligand is comprised of an antibody fused to a second monomer of a hetero-dimerizing helix protein, said first interactor domain is an antigen and said second interactor domain is a first monomer of a hetero-dimerizing helix and said second

wherein the first monomer of the hetero-dimerizing helix specifically binds to the second monomer of the hetero-dimerizing helix protein and the antibody specifically binds to the antigen-monomer of a hetero-dimerizing helix are selected from the group consisting of c-fos and c-jun.

92. (Currently amended) The polypeptide of claim 90 91, wherein the antibody is an scFv.

93-97. (Cancelled)

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- 98. (New) The polypeptide of claim 80, wherein the first interactor domain is fused through a first flexible polypeptide linker to the circularly permutated  $\beta$ -lactamase protein through the N-terminal break-point, and the second interactor domain is fused through a second flexible polypeptide linker to the circularly permutated  $\beta$ -lactamase protein through the C-terminal break-point.
- 99. (New) The polypeptide of claim 98, wherein said first polypeptide linker is 3-30 amino acids in length; and wherein said second polypeptide linker is 3-30 amino acids in length.